## CLAIMS

- A peptide which mimics a loop on the  $\gamma$ -chain that either interact with a cytokine or a  $\gamma$ -chain partner receptor chain of a heterodimeric cytokine receptor, wherein said 5 peptide consists of 5-25 amino acids and inhibits signal transduction mediated by cytokine: receptor binding of cytokines that bind to receptors that comprise a  $\gamma$ -chain.
- 2. The peptide of claim 1 wherein said peptide includes: SEQ ID NO:1 IQLYQTF; SEQ ID NO:2 IHLYQTF; SEQ ID NO:3 CLQYLV; 10 SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; SEQ 15 ID NO:17 KLSEQL or such an amino acid sequence with one or more conservative substitutions.
- 3. The peptide of claim 1 wherein said peptide includes: SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ 20 ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL or SEQ ID NO:17 KLSEQL.
- 25 4. The peptide of claim 1 wherein said peptide includes: SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 30 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; SEQ ID NO:17 KLSEQL or such an amino acid sequence with one or more conservative substitutions.

- 5. The peptide of claim 1 wherein said peptide is conformationally restricted.
- 6. The peptide of claim 1 wherein said peptide is cyclic.
- 7. The peptide of claim 1 wherein said peptide has a cysteine residue at its N terminus and a cysteine residue at its C terminus and it is cyclized by formation of a disulfide bond between said cysteine residues.
- 8. The peptide of claim 1 wherein said peptide is selected from the group consisting of: SEQ ID NO:18 CIQLYQTFC;

  10 SEQ ID NO:19 CIHLYQTFC; SEQ ID NO:20 CLQYLVC; SEQ ID NO:21 CLEHLVC; SEQ ID NO:22 CLQYLTC; SEQ ID NO:23 CLEHLTC; SEQ ID NO:24 CLQYLTQC; SEQ ID NO:25 CLEHLTQC; SEQ ID NO:26 CPIAGSSQQC; SEQ ID NO:37 CPICGSSQQC; SEQ ID NO:27 CPLCGSAQHC; SEQ ID NO:28 CPLAGSAQHC; SEQ ID NO:29 CNHEPRFLSC; SEQ ID NO:30 CDYRHKFSLC;

  15 SEQ ID NO:31 CLNLQNLC; SEQ ID NO:32 CLKLQNLC; SEQ ID NO:33 CNLSESQLC; SEQ ID NO:34 CKLSESQLC and derivative peptides thereof.
- 9. The peptide of claim 1 wherein said peptide is selected from the group consisting of: SEQ ID NO:18 CIQLYQTFC;
  20 SEQ ID NO:19 CIHLYQTFC; SEQ ID NO:20 CLQYLVC; SEQ ID NO:21 CLEHLVC; SEQ ID NO:22 CLQYLTC; SEQ ID NO:23 CLEHLTC; SEQ ID NO:24 CLQYLTQC; SEQ ID NO:25 CLEHLTQC; SEQ ID NO:26 CPIAGSSQQC; SEQ ID NO:37 CPICGSSQQC; SEQ ID NO:27 CPLCGSAQHC; SEQ ID NO:28 CPLAGSAQHC; SEQ ID NO:29 CNHEPRFLSC; SEQ ID NO:30 CDYRHKFSLC; SEQ ID NO:31 CLNLQNLC; SEQ ID NO:32 CLKLQNLC; SEQ ID NO:33 CNLSESQLC and SEQ ID NO:34 CKLSESQLC.
  - 10. The peptide of claim 1 wherein said peptide is a conformationally restricted peptide having the formula:

$$R_1 - R_2 - R_3 - R_4 - R_5$$

30 wherein:

 $R_1$  is a linking moiety;  $R_2$  is 0-10 amino acids;

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R<sub>3</sub> is SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; SEQ ID NO:17 KLSEQL or such an amino acid sequence with one or more conservative substitutions;

 $R_4$  is 0-10 amino acids; and  $R_5$  is a linking moiety.

- 11. The peptide of claim 10 wherein:  $R_1$  is cysteine; and  $R_5$  is cysteine.
- 12. The peptide of claim 11 wherein:

R<sub>3</sub> is SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; or SEQ ID NO:17 KLSEQL.

- 13. The peptide of claim 12 wherein:  $R_1$  is cysteine; and  $R_5$  is cysteine.
- 25 14. A pharmaceutical composition comprising a peptide of claim 1 and a pharmaceutically acceptable carrier or diluent.
- 15. The pharmaceutical composition of claim 14 wherein said peptide includes: SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL;

SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; SEQ ID NO:17 KLSEQL or such an amino acid sequence with one or more conservative substitutions.

- 16. The pharmaceutical composition of claim 14 wherein said peptide includes: SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL or SEQ ID NO:17 KLSEQL.
  - 17. The pharmaceutical composition of claim 14 wherein said peptide is conformationally restricted.
- 18. The pharmaceutical composition of claim 14 wherein 15 said peptide is cyclic.
  - 19. The pharmaceutical composition of claim 14 wherein said peptide has a cysteine residue at its N terminus and a cysteine residue at its C terminus and it is cyclized by formation of a disulfide bond between said cysteine residues.
- 20 20. The pharmaceutical composition of claim 14 wherein said peptide is selected from the group consisting of: SEQ ID NO:18 CIQLYQTFC; SEQ ID NO:19 CIHLYQTFC; SEQ ID NO:20 CLQYLVC; SEQ ID NO:21 CLEHLVC; SEQ ID NO:22 CLQYLTC; SEQ ID NO:23 CLEHLTC; SEQ ID NO:24 CLQYLTQC; SEQ ID NO:25 CLEHLTQC; SEQ ID NO:26 CPIAGSSQQC; SEQ ID NO:37 CPICGSSQQC; SEQ ID NO:27 CPLCGSAQHC; SEQ ID NO:28 CPLAGSAQHC; SEQ ID NO:29 CNHEPRFLSC; SEQ ID NO:30 CDYRHKFSLC; SEQ ID NO:31 CLNLQNLC; SEQ ID NO:32 CLKLQNLC; SEQ ID NO:33 CNLSESQLC; SEQ ID NO:34 CKLSESQLC and derivative peptides thereof.
- 30 21. The pharmaceutical composition of claim 14 wherein said peptide is selected from the group consisting of: SEQ ID

NO:18 CIQLYQTFC; SEQ ID NO:19 CIHLYQTFC; SEQ ID NO:20 CLQYLVC; SEQ ID NO:21 CLEHLVC; SEQ ID NO:22 CLQYLTC; SEQ ID NO:23 CLEHLTC; SEQ ID NO:24 CLQYLTQC; SEQ ID NO:25 CLEHLTQC; SEQ ID NO:26 CPIAGSSQQC; SEQ ID NO:37 CPICGSSQQC; SEQ ID NO:27 5 CPLCGSAQHC; SEQ ID NO:28 CPLAGSAQHC; SEQ ID NO:29 CNHEPRFLSC; SEQ ID NO:30 CDYRHKFSLC; SEQ ID NO:31 CLNLQNLC; SEQ ID NO:32 CLKLQNLC; SEQ ID NO:33 CNLSESQLC and SEQ ID NO:34 CKLSESQLC.

The pharmaceutical composition of claim 14 wherein 22. said peptide is a conformationally restricted peptide having 10 the formula:

$$R_1 - R_2 - R_3 - R_4 - R_5$$

wherein:

 $R_1$  is a linking moiety;

 $R_2$  is 0-10 amino acids;

15  $R_3$  is SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13

20 DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; SEQ ID NO:17 KLSEQL or such an amino acid sequence with one or more conservative substitutions;

> $R_4$  is 0-10 amino acids; and  $R_5$  is a linking moiety.

- 25 23. The pharmaceutical composition of claim 14 wherein:  $R_1$  is cysteine; and  $R_{5}$  is cysteine.
- 24. The pharmaceutical composition of claim 23 wherein: R<sub>3</sub> is SEQ ID NO:1 IQLYQTF; SEQ ID NO: 2 IHLYQTF; SEQ 30 ID NO:3 CLQYLV; SEQ ID NO:4 CLEHLV; SEQ ID NO:5 CLQYLT; SEQ ID NO:6 CLEHLT; SEQ ID NO:7 CLQYLTQ; SEQ ID NO:8 CLEHLTQ; SEQ ID NO:9 PIAGSSQQ; SEQ ID NO:36 PICGSSQQ; SEQ ID NO:10 PLCGSAQH; SEQ ID NO:11 PLAGSAQH; SEQ ID NO:12 NHEPRFLS; SEQ ID NO:13

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DYRHKFSL; SEQ ID NO:14 LNLQNL; SEQ ID NO:15 LKLQNL; SEQ ID NO:16 NLSESQL; or SEQ ID NO:17 KLSEQL.

- 25. The pharmaceutical composition of claim 24 wherein:  $R_1$  is cysteine; and  $R_5$  is cysteine.
- 26. A method of inhibiting signal transduction mediated by cytokine:receptor binding of cytokines that bind to receptors that comprise a  $\gamma$ -chain, said method comprising the step of administering an effective amount of a peptide of claim 1.
- 27. A method of inhibiting cytokine mediated cell growth, proliferation, function or activity comprising the step of contacting said cell with an peptide of claim 1.
- 28. A method of treating a patient suffering from a disease disorder or condition characterized by cytokine mediated cell growth, proliferation, function or activity comprising the step of administering to said patient a therapeutically effective amount of a peptide of claim 1.
- 29. A method of treating a patient suffering from a disease disorder or condition characterized by cytokine mediated cell growth, proliferation, function or activity comprising the step of administering to said patient a therapeutically effective amount of a peptide of claim 1, wherein said patient has lymphoma, leukemia, an allergic reaction, an autoimmune disease, graft versus host disease or rejection of a transplant or graft.
- 30. A method of preventing a condition characterized by cytokine mediated cell growth, proliferation, function or activity in a patient identified as being at risk of such a condition comprising the step of administering to said patient a prophylactically effective amount of a peptide of claim 1.

31. A method of preventing a condition characterized by cytokine mediated cell growth, proliferation, function or activity in a patient identified as being at risk of such a condition comprising the step of administering to said patient a prophylactically effective amount of a peptide of claim 1, wherein said patient is at risk of an allergic reaction, graft versus host disease or rejection of a transplant or graft.